#### AMENDMENTS TO THE DRAWINGS

The attached sheets of drawings include changes to Figure 1 and Figure 2. These sheets replace the original sheets including Figure 1 and Figure 2. In Figures 1 and 2, sequence identifier numbers (SEQ ID NO:) have been added to identify each amino acid residue sequence in the figures. No new matter is added by this amendment.

Attachment: A total of 2 Replacement Sheets containing Figure 1 (1 sheet) and Figure 2 (1 sheet)

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# Figure 2

CDRH2			
HLA-A3	N <b>V</b> G <b>SI</b> YNQK	708	(SEQ ID NO: 23)
	I <b>V</b> K <b>SI</b> TVSA	CEA	(SEQ ID NO: 24)
Pan DR	INPNN <b>V</b> G <b>SI</b>	708	(SEQ ID NO: 25)
	SI <b>V</b> K <b>SI</b> TVSA	CEA	(SEQ ID NO: 26)
HLA-DR1	<b>V</b> G <b>SI</b> YNQKF	708	(SEQ ID NO: 27)
TIEN DIXI	SI <b>V</b> K <b>SI</b> TVS		(SEQ ID NO: 28)
HLA-DR1	I <b>N</b> PNVGSI	708	(SEQ ID NO: 25)
	LATR <b>N</b> NSI	CEA	(SEQ ID NO: 29)
HLA-DR7	VGSIYN	708	(SEQ ID NO: 30)
	IVKSITV	CEA	(SEQ ID NO: 31)
CDRH3			
HLA-A3	CAR <b>GY</b> GN <b>Y</b> V	708	(SEQ ID NO: 32)
	HLF <b>GY</b> SW <b>Y</b> K	CEA	(SEQ ID NO: 33)
	NRFGY <u>S</u> WYK	NCA	(SEQ ID NO: 34)

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## Figure 1

CDRH	12							
1.	50	GINP <b>NN</b> VGSI <u>Y</u> NQKFR <u>G</u>		66	708	(SEQ ID NO: 5)		
	254	NITV <b>NN</b> SGS <u>Y</u> TCQAHNS		290	CEA	(SEQ ID NO: 15)		
	288	NITV <b>NN</b> SGSYMCQAHNS		304	NCA	(SEQ ID NO: 16)		
2.	50	GIN <b>PNN</b> V <b>GS</b> I <u>Y</u> NQKFRG		66	708	(SEQ ID NO: 5)		
	610	KIT <b>PNN</b> NG <u>TY</u> ACFVSNL		626	CEA	(SEQ ID NO: 17)		
	288	NITV <b>NN</b> S <b>GS</b> YMCQAHNS		304	NCA	(SEQ ID NO: 16)		
3.	50	<b>G</b> I <b>N</b> P <u>N</u> N <b>V</b> G <b>SI</b> YNQKFR <u>G</u>		66	708	(SEQ ID NO: 5)		
	629	GRNNSIVKSITVSASGT		645	CEA	(SEQ ID NO: 18)		
CDRH3								
1	100	<b>GY</b> GN <b>Y</b> VAY	107	708		(SEQ ID NO: 6)		
	30	<b>GY</b> SW <b>Y</b> KGE	37	CEA		(SEQ ID NO: 19)		
	59	<b>GY</b> SW <b>Y</b> KGE	72	NCA		(SEQ ID NO: 19)		
2.	100	<u>G</u> <b>Y</b> GN <b>Y</b> VAY	107	708		(SEQ ID NO: 6)		
	389	<u>S</u> YTYYRPG	396	CEA		(SEQ ID NO: 20)		
	242	<u>S</u> KAN <b>Y</b> RPG	255	NCA		(SEQ ID NO: 21)		
3.	100	$\mathtt{GYGNY} \mathbf{V} \underline{\mathbf{A}} \underline{\mathbf{Y}}$	107	708		(SEQ ID NO: 6)		
	480	EDKDA <b>VA</b> F	487	CEA		(SEQ ID NO: 22)		
	159	EDKDA <b>VA</b> F	165	NCA		(SEQ ID NO: 22)		